

SEQUENCE LISTING

10> Pharmacia & Upjohn

<120> Novel Vitamin D Receptor Related Polypeptides, Nucleic Acid Sequence Encoding the Same and Uses Thereof

<130> 10806-65

<140> US 09/143,828

<141> 1998-08-31

<160> 4

<170> PatentIn Ver. 2.0

<210> 1

<211> 2905

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: [cDNA of encoding sequence of vitamin D receptor related gamma (VDRRg)]

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<210> 2 <211> 434 <212> PRT <213> Artificial Sequence

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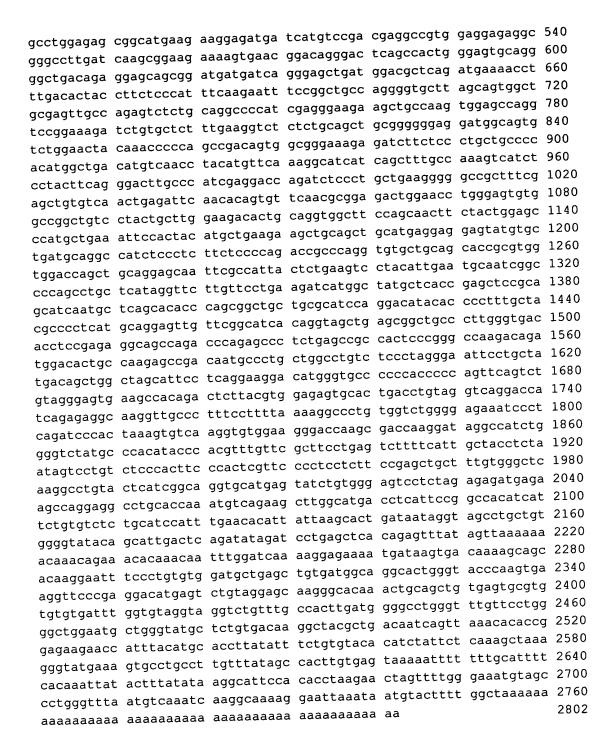
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Glu Glu Val Gly Gly Pro Gln Ile Cys Arg Val Cys Gly Asp Lys Ala 35

- Thr Gly Tyr His Phe Asn Val Met Thr Cys Glu Gly Cys Lys Gly Phe 50 60
- Phe Arg Arg Ala Met Lys Arg Asn Ala Arg Leu Arg Cys Pro Phe Arg 65 70 75 80
- Lys Gly Ala Cys Glu Ile Thr Arg Lys Thr Arg Arg Gln Cys Gln Ala 85 90 95
- Cys Arg Leu Arg Lys Cys Leu Glu Ser Gly Met Lys Lys Glu Met Ile 100 105 110
- Met Ser Asp Glu Ala Val Glu Glu Arg Arg Ala Leu Ile Lys Arg Lys
 115 120 125
- Lys Ser Glu Arg Thr Gly Thr Gln Pro Leu Gly Val Gln Gly Leu Thr 130 135 140
- Glu Glu Gln Arg Met Met Ile Arg Glu Leu Met Asp Ala Gln Met Lys
 145 150 155 160
- Thr Phe Asp Thr Thr Phe Ser His Phe Lys Asn Phe Arg Leu Pro Gly 165
- Val Leu Ser Ser Gly Cys Glu Leu Pro Glu Ser Leu Gln Ala Pro Ser 180 185 190
- Arg Glu Glu Ala Ala Lys Trp Ser Gln Val Arg Lys Asp Leu Cys Ser 195 200 205
- Leu Lys Val Ser Leu Gln Leu Arg Gly Glu Asp Gly Ser Val Trp Asn 210 215 220
- Tyr Lys Pro Pro Ala Asp Ser Gly Gly Lys Glu Ile Phe Ser Leu Leu 225 230 230 235
- Pro His Met Ala Asp Met Ser Thr Tyr Met Phe Lys Gly Ile Ile Ser 255
- Phe Ala Lys Val Ile Ser Tyr Phe Arg Asp Leu Pro Ile Glu Asp Gln 260 265 270
- Ile Ser Leu Leu Lys Gly Ala Ala Phe Glu Leu Cys Gln Leu Arg Phe 275 280 285
- Asn Thr Val Phe Asn Ala Glu Thr Gly Thr Trp Glu Cys Gly Arg Leu 290 295 300

Ser Tyr Cys Leu Glu Asp Thr Ala Gly Gly Phe Gln Gln Leu Leu 315 310 305 Glu Pro Met Leu Lys Phe His Tyr Met Leu Lys Lys Leu Gln Leu His 330 325 Glu Glu Glu Tyr Val Leu Met Gln Ala Ile Ser Leu Phe Ser Pro Asp 345 340 Arg Pro Gly Val Leu Gln His Arg Val Val Asp Gln Leu Gln Glu Gln 360 355 Phe Ala Ile Thr Leu Lys Ser Tyr Ile Glu Cys Asn Arg Pro Gln Pro 375 Ala His Arg Phe Leu Phe Leu Lys Ile Met Ala Met Leu Thr Glu Leu 395 390 Arg Ser Ile Asn Ala Gln His Thr Gln Arg Leu Leu Arg Ile Gln Asp 410 405 Ile His Pro Phe Ala Thr Pro Leu Met Gln Glu Leu Phe Gly Ile Thr 425 420 Gly Ser <210> 3 <211> 2802 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: [cDNA of encoding sequence of vitamin D receptor related gamma-2 (VDRRg-2)]

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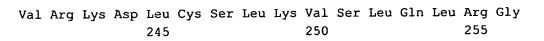
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<210> 4
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<220>

<223> Description of Artificial Sequence: [Deduced amino acid sequence of vitamin D receptor related

gamma-2 (VDRRg-2)]

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met 1	IIIT	V	aı	1111	5					10					15	
Pro	Ala	I	le	Pro 20	Leu	His	Ser	Ala	Ala 25	Ala	Glu	Leu	Ala	Ser 30	Asn	His
Pro	Arg		1y 35	Pro	Glu	Ala	Asn	Leu 40	Glu	Val	Arg	J Pro	Lys 45	Glu	Ser	Trp
Asn	His		la	Asp	Phe	Val	His		Glu	ı Asp	Thi	Glu 60	Ser	Val	Pro	Gly
Lys 65	Pro	o S	Ser	Val	Asn	Ala 70		Glu	ı Glı	ı Val	l Gl ₃	y Gly 5	Pro	Gln	Ile	Cys 80
Arg	Va:	1 (Cys	Gly	Asp 85		s Ala	a Thr	Gl:	у Ту: 9	r Hi	s Phe	. Asn	Val	Met 95	Thr
Cys	Gl	u (Gly	Cys		s Gl	y Ph	e Phe	e Ar	g Ar 5	g Al	a Met	: Lys	Arg 110	Asn	Ala
Arg	Le		Arg 115		s Pro	o Ph	e Ar	g Ly:	s Gl O	y Al	а Су	s Glu	11e	Thr	Arg	Lys
Thr	Ar 13		Arg	g Gl	n Cy	s Gl	n Al 13	а Су 5	s Ar	g Le	eu Ar	g Ly:	s Cys	s Leu	Glu	Ser
Gl ₃		ŧt	Lys	s Ly	s Gl	u Me 15		e Me	t Se	er As	sp G]	lu Al	a Val	l Glu	ı Glu	Arg 160
Arg	g Al	La	Lei	u Il	e Ly 16		g Ly	rs Ly	rs Se	er Gi	lu A: 70	rg Th	r Gl	y Thi	: Glr 175	n Pro
Le	u G	lу	۷a	1 Gl 18		y Le	eu Tl	nr Gl	Lu G.	Lu G. 35	ln A	rg Me	t Me	t Ile 190	e Arq	g Glu
Le	u M	et	As		.a Gl	ın Me	et L	ys Ti 20	nr P	he A	sp T	hr Th	r Ph 20	e Se	r Hi	s Phe
Ly		sn 10		ie Ai	cg Le	eu P	ro G 2	ly V 15	al L	eu S	er S	er Gl	Ly Cy 20	s Gl	u Le	u Pro
G1 22		er	Le	eu G	ln A		ro S 30	er A	rg G	lu G	lu A 2	Ala A 235	la Ly	/s Tr	p Se	r Gln 240



Glu Asp Gly Ser Val Trp Asn Tyr Lys Pro Pro Ala Asp Ser Gly Gly 260 265 270

Lys Glu Ile Phe Ser Leu Leu Pro His Met Ala Asp Met Ser Thr Tyr 275 280 285

Met Phe Lys Gly Ile Ile Ser Phe Ala Lys Val Ile Ser Tyr Phe Arg 290 295 300

Asp Leu Pro Ile Glu Asp Gln Ile Ser Leu Leu Lys Gly Ala Ala Phe 305 310 315 320

Glu Leu Cys Gln Leu Arg Phe Asn Thr Val Phe Asn Ala Glu Thr Gly 325 330 335

Thr Trp Glu Cys Gly Arg Leu Ser Tyr Cys Leu Glu Asp Thr Ala Gly 340 345 350

Gly Phe Gln Gln Leu Leu Glu Pro Met Leu Lys Phe His Tyr Met 355 360 365

Leu Lys Lys Leu Gln Leu His Glu Glu Glu Tyr Val Leu Met Gln Ala 370 380

Ile Ser Leu Phe Ser Pro Asp Arg Pro Gly Val Leu Gln His Arg Val 385 390 395 400

Val Asp Gln Leu Gln Glu Gln Phe Ala Ile Thr Leu Lys Ser Tyr Ile 405 410 415

Glu Cys Asn Arg Pro Gln Pro Ala His Arg Phe Leu Phe Leu Lys Ile 420 425 430

Met Ala Met Leu Thr Glu Leu Arg Ser Ile Asn Ala Gln His Thr Gln 435 440 445

Arg Leu Leu Arg Ile Gln Asp Ile His Pro Phe Ala Thr Pro Leu Met 450 455 460

Gln Glu Leu Phe Gly Ile Thr Gly Ser 465 470